



SEQUENCE LISTING

<110> SARCABAL, PATRICIA
CROUX, CHRISTIAN
SOUCAILLE, PHILIPPE

<120> METHOD FOR PREPARING 1,3-PROPANEDIOL BY A RECOMBINANT
MICRO-ORGANISM IN THE ABSENCE OF COENZYME B12 OR ONE OF
ITS PRECURSORS

<130> CHEP:004US

<140> 10/043,639

<141> 2002-01-09

<150> PCT/FR00/01981

<151> 2000-07-07

<150> FR 99/08939

<151> 1999-07-09

<160> 14

<170> PatentIn Ver. 2.1

<210> 1

<211> 2364

<212> DNA

<213> Clostridium butyricum

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caaacagaag gccagccagc aattttaaaga agagcattgg cattgaaaca cataactgaa 180
aatatcccta taacaattag agatcaagaa cttatagtgg gaagtttaac taaagaacca 240
aggccttcac aagtatttcc tgagtttct aataagtgg tacaagatga attggataga 300
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ggcgttaggac atgtatctgt agattatgga aaagttttaa gggttggatt taatgggatt 540
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<211> 915
<212> DNA
<213> Clostridium butyricum
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aatccagaat cccaagatataaacctcaa gtaatgttta ataaaaattt atgtacaaaa 180
tgtggaaagat gtaaatctca atgtaaaagt gcaggttattt atatgaattt agaatatagg 240
atagataaaa gcaaatgtac agagtgtaca aaatgtgttgg ataaattgcctt aagcggggca 300
cttggatttgg aaggaaggaa ttacagtgtt gaagacgtt taaaggaatt gaaaaaaagat 360
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agtaacgaaa taatattaca aaacattttttaa ttaagtgtatg aatttagctaa agaaataata 660
atcagaattc ctgtataga aggatttaat gcagatttac aaagtgatgg agcaatagct 720
caattttcaa aatcatttacaa aatctttaaa agaataatgttccatc ttcttccatc ccataattat 780
ggagaaaaata agtatcaagc aattggaaaga gatgttctt tggaaagaact aaaatcacct 840
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<213> *Clostridium butyricum*

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28

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<210> 4
<211> 1158
<212> DNA
<213> Clostridium butyricum
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<210> 5
<211> 4963
<212> DNA
<213> *Clostridium butyricum*

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caattgtatt agtttaact ttagataaaa caaacaaaaa tgttattatt agccaagaaa 180
atactgtac aaaagaaaaaag agaaaaacat agcaaaagag tccaatatt aagcaataaa 240
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ggaggagtaa aatgataag taaaggattt agtacccaaa cagaaaagaat aaatattta 360
aaggctcaa tattaaatgc taaaccatgt gttgaatcag aaagagcaat attaataaca 420
gaatcattt aacaaacaga aggccagcca gcaattttaa gaagagcatt ggcattgaaa 480
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gaattggata gattaaataa gagaactgga gatgcattcc aaatttaaga agaaagtaaa 660
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<210> 6
<211> 783
<212> PRT

<213> Clostridium butyricum

<400> 6

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Leu Ile Thr Glu Ser Phe Lys Gln Thr Gly Gln Pro Ala Ile Leu Arg
35 40 45

Arg Ala Leu Ala Leu Lys His Ile Leu Glu Asn Ile Pro Ile Thr Ile
50 55 60

Arg Asp Gln Glu Leu Ile Val Gly Ser Leu Thr Lys Glu Pro Arg Ser
65 70 75 80

Ser Gln Val Phe Pro Glu Phe Ser Asn Lys Trp Leu Gln Asp Glu Leu
85 90 95

Asp Arg Leu Asn Lys Arg Thr Gly Asp Ala Phe Gln Ile Ser Glu Glu
100 105 110

Ser Lys Glu Lys Leu Lys Asp Val Phe Glu Tyr Trp Asn Gly Lys Thr
115 120 125

Thr Ser Glu Leu Ala Thr Ser Tyr Met Thr Glu Glu Thr Arg Glu Ala
130 135 140

Val Asn Cys Glu Val Phe Thr Val Gly Asn Tyr Tyr Tyr Asn Gly Val
145 150 155 160

Gly His Val Ser Val Asp Tyr Lys Val Leu Arg Val Gly Phe Asn Gly
165 170 175

Ile Ile Asn Glu Ala Lys Glu Gln Leu Glu Lys Asn Arg Ser Asp Pro
180 185 190

Asp Phe Ile Lys Lys Glu Lys Phe Leu Asn Ser Val Ile Ile Ser Cys
195 200 205

Glu Ala Ala Ile Thr Tyr Val Asn Arg Tyr Ala Lys Lys Ala Lys Glu
210 215 220

Ile Ala Asp Asn Thr Ser Asp Ala Lys Arg Lys Ala Glu Leu Asn Glu
225 230 235 240

Ile Ala Lys Ile Cys Ser Lys Val Ser Gly Glu Gly Ala Lys Ser Phe
245 250 255

Tyr Glu Ala Cys Gln Leu Phe Trp Phe Ile His Ala Ile Ile Asn Ile
260 265 270

Glu Ser Asn Gly His Ser Ile Ser Pro Ala Arg Phe Asp Gln Tyr Met
275 280 285

Tyr Pro Tyr Tyr Glu Asn Asp Lys Asn Ile Thr Asp Lys Phe Ala Gln
290 295 300

Glu Leu Ile Asp Cys Ile Trp Ile Lys Leu Asn Asp Ile Asn Lys Val
305 310 315 320

Arg Asp Glu Ile Ser Thr Lys His Phe Gly Gly Tyr Pro Met Tyr Gln
325 330 335

Lys Leu Ile Val Gly Gly Gln Asn Ser Glu Gly Lys Asp Ala Thr Asn
340 345 350

Lys Val Ser Tyr Met Ala Leu Glu Ala Ala Val His Val Lys Leu Pro
355 360 365

Gln Pro Ser Leu Ser Val Arg Ile Trp Asn Lys Thr Pro Asp Glu Phe
370 375 380

Leu Leu Arg Ala Ala Glu Leu Thr Arg Glu Gly Leu Gly Leu Pro Ala
385 390 395 400

Tyr Tyr Asn Asp Glu Val Ile Ile Pro Ala Leu Val Ser Arg Gly Leu
405 410 415

Thr Leu Glu Asp Ala Arg Asp Tyr Gly Ile Ile Gly Cys Val Glu Pro
420 425 430

Gln Lys Pro Gly Lys Thr Glu Gly Trp His Asp Ser Ala Phe Phe Asn
435 440 445

Leu Ala Arg Ile Val Glu Leu Thr Ile Asn Ser Gly Phe Asp Lys Asn
450 455 460

Lys Gln Ile Gly Pro Lys Thr Gln Asn Phe Glu Glu Met Lys Ser Phe
465 470 475 480

Asp Glu Phe Met Lys Ala Tyr Lys Ala Gln Met Glu Tyr Phe Val Lys
485 490 495

His Met Cys Cys Ala Asp Asn Cys Ile Asp Ile Ala His Ala Glu Arg
500 505 510

Ala Pro Leu Pro Phe Leu Ser Ser Met Val Asp Asn Cys Ile Gly Lys
515 520 525

Gly Lys Ser Leu Gln Asp Gly Gly Ala Glu Tyr Asn Phe Ser Gly Pro
530 535 540

Gln Gly Val Gly Val Ala Asn Ile Gly Asp Ser Leu Val Ala Val Lys
545 550 555 560

Lys Ile Val Phe Asp Glu Asn Lys Ile Thr Pro Ser Glu Leu Lys Lys
565 570 575

Thr Leu Asn Asn Asp Phe Lys Asn Ser Glu Glu Ile Gln Ala Leu Leu
580 585 590

Lys Asn Ala Pro Lys Phe Gly Asn Asp Ile Asp Glu Val Asn Leu
595 600 605

Ala Arg Glu Gly Ala Leu Val Tyr Cys Arg Glu Val Asn Lys Tyr Thr
610 615 620

Asn Pro Arg Gly Gly Asn Phe Gln Pro Gly Leu Tyr Pro Ser Ser Ile
625 630 635 640

Asn Val Tyr Phe Gly Ser Leu Thr Gly Ala Thr Pro Asp Gly Arg Lys
645 650 655

Ser Gly Gln Pro Leu Ala Asp Gly Val Ser Pro Ser Arg Gly Cys Asp
660 665 670

Val Ser Gly Pro Thr Ala Ala Cys Asn Ser Val Ser Lys Leu Asp His
675 680 685

Phe Ile Ala Ser Asn Gly Thr Leu Phe Asn Gln Lys Phe His Pro Ser
690 695 700

Ala Leu Lys Gly Asp Asn Gly Leu Met Asn Leu Ser Ser Leu Ile Arg
705 710 715 720

Ser Tyr Phe Asp Gln Lys Gly Phe His Val Gln Phe Asn Val Ile Asp
725 730 735

Lys Lys Ile Leu Leu Ala Ala Gln Lys Asn Pro Glu Lys Tyr Gln Asp
740 745 750

Leu Ile Val Arg Val Ala Gly Tyr Ser Ala Gln Phe Ile Ser Leu Asp
755 760 765

Lys Ser Ile Gln Asn Asp Ile Ile Ala Arg Thr Glu His Val Met
770 775 780

<210> 7
<211> 304
<212> PRT
<213> Clostridium butyricum

<400> 7
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Leu His Asp Gly Pro Gly Ile Arg Thr Ile Val Phe Phe Lys Gly Cys
20 25 30

Ser Met Ser Cys Leu Trp Cys Ser Asn Pro Glu Ser Gln Asp Ile Lys
35 40 45

Pro Gln Val Met Phe Asn Lys Asn Leu Cys Thr Lys Cys Gly Arg Cys
50 55 60

Lys Ser Gln Cys Lys Ser Ala Gly Ile Asp Met Asn Ser Glu Tyr Arg
65 70 75 80

Ile	Asp	Lys	Ser	Lys	Cys	Thr	Glu	Cys	Thr	Lys	Cys	Val	Asp	Asn	Cys
85														95	
Leu	Ser	Gly	Ala	Leu	Val	Ile	Glu	Gly	Arg	Asn	Tyr	Ser	Val	Glu	Asp
100														110	
Val	Ile	Lys	Glu	Leu	Lys	Asp	Ser	Val	Gln	Tyr	Arg	Arg	Ser	Asn	
115														125	
Gly	Gly	Ile	Thr	Leu	Ser	Gly	Gly	Glu	Val	Leu	Leu	Gln	Pro	Asp	Phe
130														140	
Ala	Val	Glu	Leu	Leu	Lys	Glu	Cys	Lys	Ser	Tyr	Gly	Trp	His	Thr	Ala
145														155	160
Ile	Glu	Thr	Ala	Met	Tyr	Val	Asn	Ser	Glu	Ser	Val	Lys	Lys	Val	Ile
165														175	
Pro	Tyr	Ile	Asp	Leu	Ala	Met	Ile	Asp	Ile	Lys	Ser	Met	Asn	Asp	Glu
180														190	
Ile	His	Arg	Lys	Phe	Thr	Gly	Val	Ser	Asn	Glu	Ile	Ile	Leu	Gln	Asn
195														205	
Ile	Lys	Leu	Ser	Asp	Glu	Leu	Ala	Lys	Glu	Ile	Ile	Arg	Ile	Pro	
210														220	
Val	Ile	Glu	Gly	Phe	Asn	Ala	Asp	Leu	Gln	Ser	Ile	Gly	Ala	Ile	Ala
225														235	240
Gln	Phe	Ser	Lys	Ser	Leu	Thr	Asn	Leu	Lys	Arg	Ile	Asp	Leu	Leu	Pro
245														255	
Tyr	His	Asn	Tyr	Gly	Glu	Asn	Lys	Tyr	Gln	Ala	Ile	Gly	Arg	Glu	Tyr
260														270	
Ser	Leu	Lys	Glu	Leu	Lys	Ser	Pro	Ser	Lys	Asp	Lys	Met	Glu	Arg	Leu
275														285	
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290														300	

<210> 8
 <211> 385
 <212> PRT
 <213> Clostridium butyricum

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20

25

30

Lys Ala Leu Ile Val Thr Asp Lys Phe Leu Lys Asp Met Glu Gly Gly
35 40 45

Ala Val Glu Leu Thr Val Lys Tyr Leu Lys Glu Ala Gly Leu Asp Val
50 55 60

Val Tyr Tyr Asp Gly Val Glu Pro Asn Pro Lys Asp Val Asn Val Ile
65 70 75 80

Glu Gly Leu Lys Ile Phe Lys Glu Glu Asn Cys Asp Met Ile Val Thr
85 90 95

Val Gly Gly Ser Ser His Asp Cys Gly Lys Gly Ile Gly Ile Ala
100 105 110

Ala Thr His Glu Gly Asp Leu Tyr Asp Tyr Ala Gly Ile Glu Thr Leu
115 120 125

Val Asn Pro Leu Pro Pro Ile Val Ala Val Asn Thr Thr Ala Gly Thr
130 135 140

Ala Ser Glu Leu Thr Arg His Cys Val Leu Thr Asn Thr Lys Lys Lys
145 150 155 160

Ile Lys Phe Val Ile Val Ser Trp Arg Asn Leu Pro Leu Val Ser Ile
165 170 175

Asn Asp Pro Met Leu Met Val Lys Lys Pro Ala Gly Leu Thr Ala Ala
180 185 190

Thr Gly Met Asp Ala Leu Thr His Ala Ile Glu Ala Tyr Val Ser Lys
195 200 205

Asp Ala Asn Pro Val Thr Asp Ala Ser Ala Ile Gln Ala Ile Lys Leu
210 215 220

Ile Ser Gln Asn Leu Arg Gln Ala Val Ala Leu Gly Glu Asn Leu Glu
225 230 235 240

Ala Arg Glu Asn Met Ala Tyr Ala Ser Leu Leu Ala Gly Met Ala Phe
245 250 255

Asn Asn Ala Asn Leu Gly Tyr Val His Ala Met Ala His Gln Leu Gly
260 265 270

Gly Leu Tyr Asp Met Ala His Gly Val Ala Asn Ala Met Leu Leu Pro
275 280 285

His Val Glu Arg Tyr Asn Met Leu Ser Asn Pro Lys Lys Phe Ala Asp
290 295 300

Ile Ala Glu Phe Met Gly Glu Asn Ile Ser Gly Leu Ser Val Met Glu
305 310 315 320

Ala Ala Glu Lys Ala Ile Asn Ala Met Phe Arg Leu Ser Glu Asp Val

325 330 335
Gly Ile Pro Lys Ser Leu Lys Glu Met Gly Val Lys Gln Glu Asp Phe
340 345 350

Glu His Met Ala Glu Leu Ala Leu Leu Asp Gly Asn Ala Phe Ser Asn
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Pro Arg Lys Gly Asn Ala Lys Asp Ile Ile Asn Ile Phe Lys Ala Ala
370 375 380

Tyr
385

<210> 9
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<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Primer

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<210> 10
<211> 40
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Primer

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<210> 11
<211> 34
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Primer

<400> 11
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<210> 12
<211> 34
<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
Primer

<400> 12

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34

<210> 13

<211> 31

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
Primer

<400> 13

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31

<210> 14

<211> 36

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
Primer

<400> 14

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36